		-		က
		NO:		NO:
		ID		ID
		SEQ		SEQ
1294	_	GTAGGAAAA		GTAGGAAAA
1284	-	CAAATTATAAACATGTGGCAGAAAGTAGGAAAA		CAAATTAAAAACTTTTTAAAGAAAGTAGGAAAA
1274		TATAAACAT		TAAAAACTT
1264		CAAAT	<u>=</u>	CAAAT
.		gp120 423-432		rv_85283

Figure 2

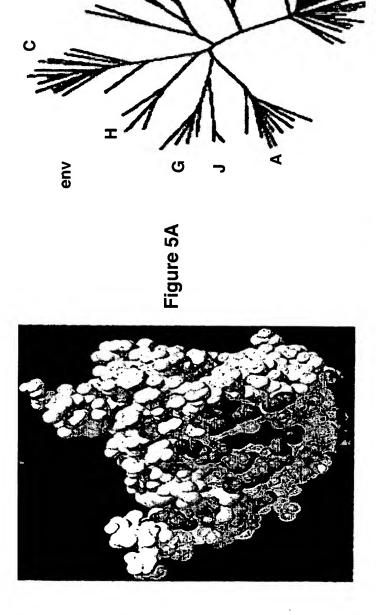


Figure 5C

NAME	SEQUENCE	SEO ID	NAME	SEOUENCE	SEO ID
gp120	KQIINM-WQR-VGKAMY	48	G.BE.DRBCL	KOIINM-WOR-VGKAMY	59
(421-436)				1	
A.GB.MA245	KQIINM-WQR-VGKAMY	49	H.BE.VI991	KQIINM-WQR-VGKAMY	09
A.GB.MC108	KQIINM-WQR-VGKAMY	50		•	
B.AU.MBC18	KQIINM-WQR-VGKAMY	51	J.SE.SE9173	KQIINM-WQR-VGKAMY	61
B.AU.MBC200	KQIINM-WQR-VGKAMY	52			
C.BI.BU910112	KQIINM-WQR-VGKAMY	53	K.CD.EQTB11C	KQIINM-WQR-VGKAMY	62
C.BI.BU910316	KQI INM-WQR-VGKAMY	54			
D.CD.84ZR085	KQIINM-WQR-VGKAMY	55	N.CM.YBF30	KQIINM-WQR-VGKAMY	63
D.CD.JY1	KQIINM-WQR-VGKAMY	56			
F.BR.BZ126A	KQIINM-WQR-VGKAMY	57	O.CMANT70C	KQIINM-WQR-VGKAMY	64
F1.BE.VI850	KQI INM-WQR-VGKAMY	58			

Figure 5B

Replacement Sheet

scFv JL413 VL domain

PRO SER PRO THR THR LYS SER GLY VAL PRO GLN GLYTHR GLN SER PRO SER SER VAL SER ALA SER VAL GLY ASP ARG VAL LEU GLN GLN GLYSER GLY SER PHE TYR THR TRP SER LEU GLN ILE THR TRP LEU ALA GLY LEU LEU VAL SER THR GLY ASN PHE TYR CYS GLN GLN ALA GLY ALA ASP 46 SEQ ID NO: ILE TYR THR ILE SER GLN GLY GLY ASN LEU LEU SER TYR GLY ILE LYS ARG ALA THR SER HIS MET THR CYS ARG SER ALA GLXALA SER LYS VAL GLU ASP VAL VAL GLY LYS GLU ASP ARG PHE ILE 101

scFv JL413 VL domain

PRO TYR GLYLEU ARG GLU SER GLY PRO GLY LEU VAL LYS PRO SER GLU THR LEU SER LEU LEU GLN SER GLN ASN ARG THR PHE GLYILE GLN TRP HIS CYS SER TRP SER VAL ASP THR SER LYS ASN TYR SER GLY CXS TRP SER TYR GLY ARG THR ALA MET TYR TYR 47 SER ILE . ON TYR SER SER SEQ ID ALA ASP ILE SER ILE THR GLY PHE TRP VAL ALA LYS SER ARG VAL SER GLXTHR GLY LEU GLU THR VAL SER VAL SER SER THR VAL GLN VAL ASN LEU GLY LYS SER LEU VAL CXS SER LEU THR PRO PRO

Figure 10A

scFv JL427 VL domain

ILE PHE ARG VAL TYR ARG ASN ASP GLN ARG PRO LEU GLY VAL PRO HIS THR TRP GLY LEU GLN SER VAL LEU THR GLN PRO PRO SER VAL SER GLY ALA PRO GLY GLN ARG VAL TYR GLN GLY SER SER LEU SER SER SER SER ASN PHE GLY LEU ASN TYR VAL TYR TRP ILE SER LEU ALA 44 SEQ ID NO: TYR ASP ASN THR SER ALA TYR CYS GLN SER PHE GLY GLY TYR GLN LEU TYR VAL LEU GLY LEU ILE SER GLY CDR1 ALA PRO LYS LEU GLU ALA ASP TYR SER LYS GLY SER SER CYS SER GLY PRO GLY THR ALA ARG PHE SER GLU ASP 101

scFv JL427 VL domain

TYR GLY MET ASP ILE TRP GLY GLN GLY THR THR VAL THR VAL SER SEQ ID NO: ALA TYR TYR LEU LEU 09 GLN VAL GLN LEU GLN GLN SER GLY GLY GLY LEU VAL GLN PRO GLY ARG SER LEU ARG ASN SER LEU ARG ALA GLU ASP THR ALA VAL TYR TYR CYS ALA ARG GLY ASP SER VAL LYS GLY ARG PHE THR ILE SER ARG ASP ASN SER LYS ASN THR LEU ARG THR VAL HIS TRP SER HIS GLY ARG SER GLY GLY MET TYR SER TYR ILE SER GLY LYS GLY LEU GLU TRP VAL SER THR PHE GLY PHE SER ASN CYS ALA ALA ILE ASN LEU GLN SER PRO ALA

Figure 10B

THR	PRO	SER	80 PRO	GLN
VAL	GLN	PRO	GLN	GLY
ARG	GLN	VAL PRO	reu	PHE
ASP		GLY	SER	THR
GLY	TRP TYR GLN	SER	ILE SER SER LEU	ARG
VAL	TRP	GLN	ILE	PRO
SER VAL GLY ASP ARG	ASN	SER LEU GLN SER GLY	THR	TYR SER ILE PRO ARG THR PHE
SER ALA	TYR LEU	SER	LEU	SER
SER	TYR	SER	THR	TYR
LEU		ILE TYR ALA ALA SER		SER
SER SER LEU	SER SER	ALA	ASP PHE	CYS GLN GLN SER
SER	ILE	TYR	THR	GEN
PRO	SER	ILE	GLY	CYS
SER	GEN	LEU	SER	PHE
GLN	THR CYS ARG ALA SER GLI	LEU	SER GLY SER GLY SER	ASP PHE ALA THR RYR PHE
THR	ALA	LYS	SER	THR
MET	ARG	LYS ALA PRO LYS LEU	GLY	ALA
GLN	CYS	ALA	SER	PHE
1 ASP ILE GLN MET THR GLN SE		LYS	PHE	
1 ASP	21 VAL	41 GLY	61 ARG	81 GLU

Light chain SK18 VL domain

Figure 10C

SEQ ID NO: 43

THR LYS VAL GLU ILE LYS

GLY101